

Supplementary Table 1. Multivariate regression analysis of the factors associated with tumor necrosis factor receptor 1 or tumor necrosis factor receptor 2 in healthy subjects.

<b>TNFR1</b>					<b>TNFR2</b>				
Variables	Estimate	SE	t value	P	Variables	Estimate	SE	t value	P
eGFR-CysC	-0.0019	0.0002	-8.859	<0.0001	eGFR-CysC	-0.0018	0.0002	-7.782	<0.0001
eGFR-Cr	-0.00007	0.0003	-0.229	0.82	eGFR-Cr	-0.0001	0.0004	-0.290	0.77
					Age	-0.0005	0.0004	-1.047	0.30
					Systolic BP	0.0006	0.0003	1.968	0.0498

Abbreviations: BP, blood pressure; Cr, creatinine; CysC, cystatin C; eGFR, estimated glomerular filtration rate; SE, standard error; TNF, tumor necrosis factor; TNFR, TNF receptor.

Supplementary Table 2. Multivariate regression analysis of the factors associated with tumor necrosis factor receptor 1 or tumor necrosis factor receptor 2 in patients with type 2 diabetes.

<b>TNFR1</b>					<b>TNFR2</b>				
Variables	Estimate	SE	t value	P	Variables	Estimate	SE	t value	P
eGFR-CysC	-0.0016	0.0003	-5.69	<0.0001	eGFR-CysC	-0.0017	0.0003	-6.01	<0.0001
eGFR-Cr	-0.0006	0.0004	-1.48	0.14	eGFR-Cr	-0.0003	0.0004	-0.66	0.51
Age	0.0004	0.0007	0.56	0.57	Age	0.0011	0.0007	1.48	0.14
BMI	0.0013	0.0013	1.06	0.29	HbA1c	0.0122	0.00535	2.27	0.02
HbA1c	0.0191	0.0049	3.9	0.0001	Uric acid	0.0128	0.0052	2.45	0.01
Uric acid	0.0175	0.0048	3.52	0.0003	HDL-C	-0.0023	0.0005	-4.4	<0.0001
Non-HDL-C	-0.0004	0.0002	-2.41	0.02	Non-HDL-C	-0.0005	0.0002	-2.8	0.006
UACR	0.0482	0.0086	5.62	<0.0001	UACR	0.0345	0.0091	3.78	0.0002

Abbreviations used in this table are the same as in Supplementary Table 1.

Abbreviations: BMI, body mass index; HbA1c, glycated hemoglobin; HDL-C, high-density-lipoprotein cholesterol; UACR, urinary albumin-to-creatinine ratio.

Supplementary Table 3. Spearman's correlation between tumor necrosis factor receptors and clinical parameters in healthy subjects and patients with type 2 diabetes.

Variables	Healthy subjects and diabetes (n = 705)	
	TNFR1	TNFR2
Age	0.37 <sup>***</sup>	0.45 <sup>***</sup>
BMI	0.34 <sup>***</sup>	0.33 <sup>***</sup>
Systolic BP	0.35 <sup>***</sup>	0.39 <sup>***</sup>
Diastolic BP	0.25 <sup>***</sup>	0.30 <sup>***</sup>
HbA1c	0.52 <sup>***</sup>	-0.57 <sup>***</sup>
HDL-C	-0.33 <sup>***</sup>	-0.35 <sup>***</sup>
Non-HDL-C	-0.07	-0.09 <sup>*</sup>
Uric acid	0.11 <sup>**</sup>	0.09 <sup>*</sup>
UACR	0.42 <sup>**</sup>	0.44 <sup>***</sup>
GFR-Cr	-0.10 <sup>**</sup>	-0.10 <sup>**</sup>
GFR-CysC	-0.43 <sup>***</sup>	-0.43 <sup>***</sup>
TNFR2	0.81 <sup>***</sup>	—

\* p<0.05, \*\* p<0.01, \*\*\* p<0.001

Abbreviations used in this table are the same as in Supplementary Table 1 and Table 2.

Supplementary Table 4. Multivariate stepwise regression analysis of the factors associated with tumor necrosis factor 1 or tumor necrosis factor 2 in healthy subjects and patients with type 2 diabetes.

<b>TNFR1</b>					<b>TNFR2</b>				
Variables	Estimate	SE	t value	P	Variables	Estimate	SE	t value	P
BMI	0.0013	0.0009	1.466	0.14	Age	0.0010	0.0004	2.833	0.005
HbA1c	0.0025	0.0034	7.422	<0.0001	HbA1c	0.0294	0.0037	7.850	<0.0001
HDL-C	-0.0008	0.00023	-3.520	0.0005	HDL-C	-0.0008	0.000247	-3.381	<0.0001
Non-HDL	-0.0005	0.0001	-4.645	<0.0001	Non-HDL	-0.0006	0.0001	-5.535	<0.0001
Uric acid	0.0094	0.0029	3.210	<0.0001	Uric acid	0.0104	0.0031	3.379	0.0008
eGFR-CysC	-0.0019	0.0002	-11.622	<0.0001	SBP	0.0004	0.0002	1.782	0.08
eGFR-Cr	-0.0005	0.0002	-2.095	0.04	eGFR-CysC	-0.0019	0.0002	-9.894	<0.0001
ACR	0.0441	0.0068	6.444	<0.0001	ACR	0.0383	0.00752	5.087	<0.0001

Abbreviations used in this table are the same as in Supplementary Table 1 and Table 2.

Supplementary Table 5. TNFR levels in patients with diabetes.

Author	Type of diabetes	n	Trial, study, and/or ethnicity	Age (yrs)	HbA1c (%)	eGFR-Cr or mGFR (mL/min/1.73 m <sup>2</sup> )	eGFR-CysC (mL/min/1.73 m <sup>2</sup> )	Median UACR or UAER**	TNFR1 (pg/mL)	TNFR2 (pg/mL)	Vendor (ELISA Cat. No.)***	References
Gohda T, et al.	2	292	Japanese	57±11	7.5±1.2	85±19	97±23	18 (7, 59) mg/gCr	1357 (1133, 1646)	2904 (2505, 3391)	R&D Systems (DRT100, DRT200)	Present study
Kamei N, et al. [12]	2	594	Japanese	55±13	7.3±1.2	69 (56, 84)	—	22 (9, 123) mg/gCr	1562 (1263, 2016)	3339 (2717, 4297)	R&D Systems (DRT100, DRT200)	Sci Rep. 8: 15302, 2018.
Murakoshi M, et al. [10]	2	201	Japanese	67±10	7.2±1.0	73±25	—	23 (8, 142) mg/gCr	1412 (1153, 1902)	Not measured	R&D Systems (DRT100)	Front Endocrinol. 13: 849457, 2022.
Saulnier PJ, et al. [28]	2	1135	SURDIAGENE study, French	64±11	7.8±1.5	76±21	—	23 (9, 92) mg/gCr	1818 (1544, 2231)	Not measured	EKF Diagnostics (BIO94TNFR1)	Diabetes Care. 40: 367-374, 2017.
Oshima M, et al. [29]	2	330	ADVANCE and ADVANCE-ON studies	68±7	7.6±1.6	67±16	—	15 (7, 37) mg/gCr	1429 (1150, 1760)	Not measured	R&D Systems (DRT100)	KI Report. 6: 284-295, 2021.
Pavkov ME, et al. [24]	2	83	Pima Indians	46±10	9.2 (7.6, 11.2)	119 (94, 155)*	—	26 (12, 127) mg/gCr	1500 (1205, 1960)	3283 (2670, 4151)	R&D Systems (DRT100, DRT200)	Kidney Int. 89: 226-234, 2016.
Pavkov ME, et al. [30]	2	193	Pima Indians	46 (39, 53)	9.6 (7.7, 11.1)	120 (88, 149)*	—	72 (19, 493) mg/gCr	2833 (2081, 4092)	4835 (3875, 6997)	R&D Systems (DRT100, DRT200)	Kidney Int. 87: 812-819, 2015.
Gohda T, et al. [6]	1	275	First Joslin Kidney Study, Caucasians	31±8	8.9±1.5	—	133±30	56 (37, 101) µg/min**	1345 (1156, 1598)	2161 (1732, 2673)	R&D Systems (DRT100, DRT200)	J Am Soc Nephrol. 23: 516-524, 2012.
Gohda T, et al. [6]	1	353	Second Joslin Kidney Study, Caucasians	39±12	8.4±1.3	—	129±30	41 (24, 79) µg/min**	1382 (1180, 1709)	2230 (1869, 2695)	R&D Systems (DRT100, DRT200)	J Am Soc Nephrol. 23: 516-524, 2012.

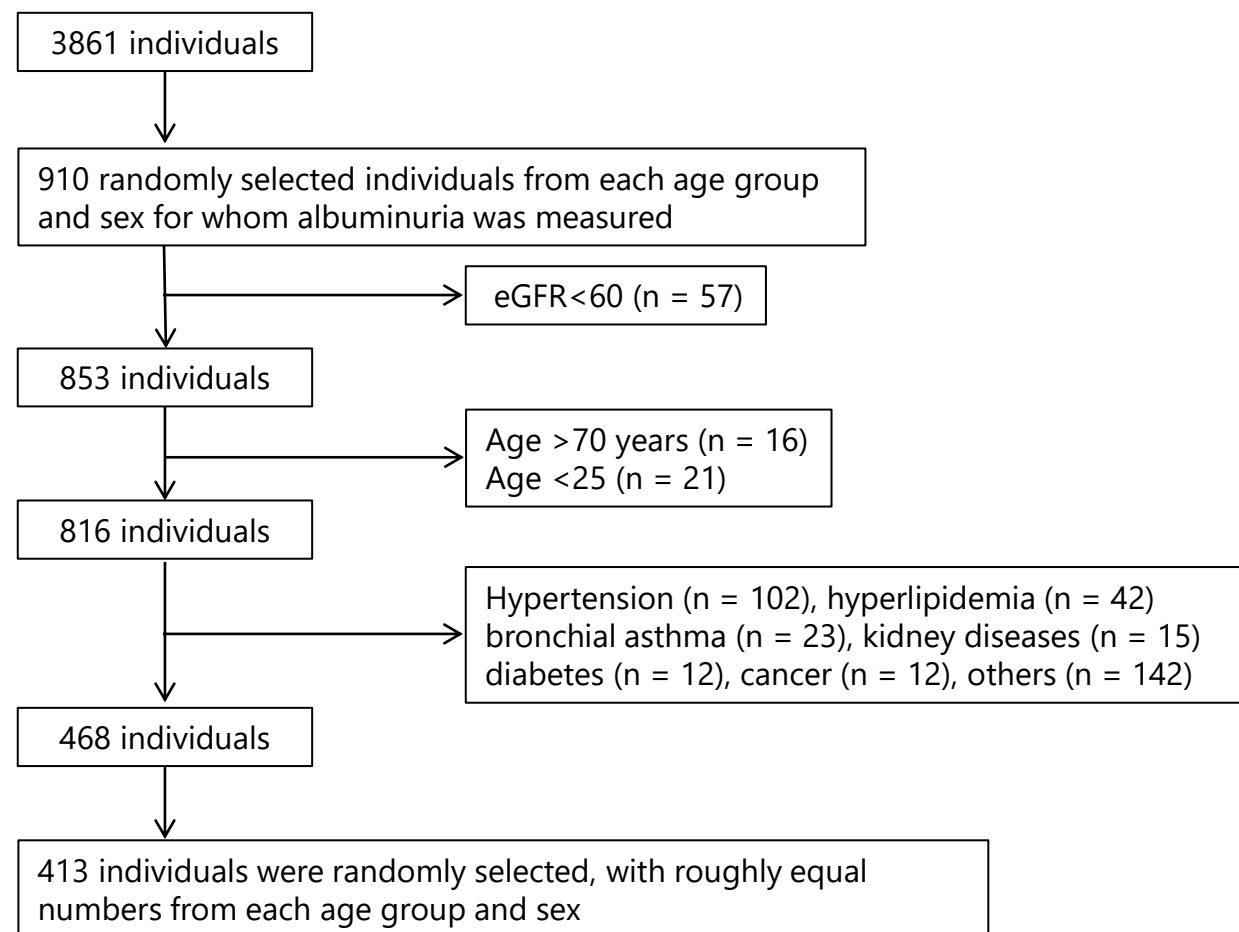
Abbreviations used in this table are the same as in Supplementary table 1.

Data are presented as mean ± standard deviation (SD) or median (quartiles). n, number of patients; \*mGFR, measured GFR (urinary clearance of nonradioactive iothalamate); \*\*AER (mg/min), albumin excretion rate

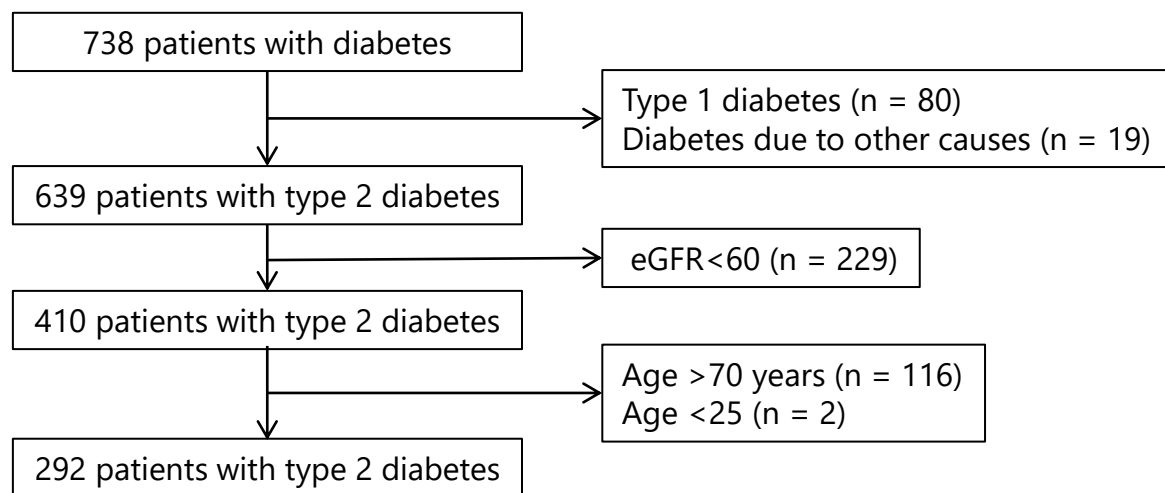
\*\*\*DRT100 (R&D Systems) and BIO94TNFR1 (EKF Diagnostics) are the same ELISA kit, although the vendor is different.

# Supplementary Figure 1

## Healthy subjects



## Type 2 diabetes



**Figure legend**

**Supplementary Figure 1.** Flowchart of inclusion and exclusion of subjects in this study.